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For all sequences using "Artificial sequence", for numeric identifier
<213>, a mandatory feature is required to explain the source of the
genetic material. The feature consists of <220>, which remains blank,
and <223>, which states the source of the genetic material. To explain
the source, if the sequence is put together from several organisms,
please list those organisms. If the sequence is made in the laboratory,
please indicate that the sequence is synthesized. These errors appear in
other sequences in the sequence listing. Please make all necessary
changes.

* *

Application No: 10576988   Version No: 1.0

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Output Set:

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Finished:	2008-10-08 19:20:51.787
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Total Errors:	1
No. of SeqIDs Defined:	10
Actual SeqID Count:	10

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W 213	Artificial or Unknown found in <213> in SEQ ID (10)

SEQUENCE LISTING

<110>   Rodriguez Aguirre et al.

<120>   EMPTY CAPSIDS (VLPs(-VP4)) OF THE INFECTIOUS BURSAL DISEASE VIRUS
(IBDV), OBTAINMENT PROCESS AND APPLICATIONS

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Leu Met Pro Thr Thr Gly Pro Ala Ser Ile Pro Asp Asp Thr Leu Glu
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Lys His Thr Leu Arg Ser Glu Thr Ser Thr Tyr Asn Leu Thr Val Gly
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Cys Arg Leu Val Ser Arg Ser Leu Thr Val Arg Ser Ser Thr Leu Pro
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Ala Thr Ala Asn Ile Asn Asp Lys Ile Gly Asn Val Leu Val Gly Glu
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agg ctt ggt gac ccc att ccc gca ata ggg ctt gac cca aaa atg gta        872
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Val Gly Asp Thr Gly Ser Gly Leu Ile Val Phe Phe Pro Gly Phe Pro
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Asn Tyr Cys Arg Leu Val Ser Arg Ser Leu Thr Val Arg Ser Ser Thr
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Phe Gln Gly Ser Leu Ser Glu Leu Thr Asp Val Ser Tyr Asn Gly Leu
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Met Ser Ala Thr Ala Asn Ile Asn Asp Lys Ile Gly Asn Val Leu Val
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Gly Glu Gly Val Thr Val Leu Ser Leu Pro Thr Ser Tyr Asp Leu Gly
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